

#	Position	Sequence context	Structural context	Local structure visualization
1	133	GGGUACUUCCGGGCA CGCAC AGACC CCCGG GGUGC ACCCU GCACC	N/A	N/A
2	324	UGCGC CCUGC GGGGG AGACC UGACU UUUGA CCCCG GCCCC CGGGA	N/A	N/A
3	346	ACUUU UGACC CCGGC CCCCG GGACC UGGUG GAGCC CUGGG UGGUG	N/A	N/A
4	508	GAGGU ACCUC CUGCA GUGUC UGACA UCCGG GUGAC GCGGU CCUCA	N/A	N/A
5	589	GCACC CAGUG GGGCU GUGCU GGACU ACGAG GUCAA AUACC AUGAG	N/A	N/A
6	919	GGGAG AGAAC CAGAA UAUUC GGACA AACAC GGACA GUAUC UCAUC	N/A	N/A
7	929	CAGAA UAUUC GGACA AACAC GGACA GUAUC UCAUC GGACA UGGUA	N/A	N/A
8	944	AACAC GGACA GUAUC UCAUC GGACA UGGUA CUAAG GUCUA CAUCG	N/A	N/A
9	1297	UUCAU GGAGA ACGGC GCCCU GGACU CCUUC CUGCG GCUAA ACGAC	N/A	N/A
10	1322	CCUUC CUGCG GCUAA ACGAC GGACA GUUCA CAGUC AUCCA GCUCG	N/A	N/A
11	1465	AACCU CGUCU GCAAA GUGUC UGACU UUGGC CUUUC CCGAU UCCUG	N/A	N/A
12	1495	CUUUC CCGAU UCCUG GAGGA GAACU CUUCC GAUCC CACCU ACACG	N/A	N/A

13	1551	AGGAA AGAUU CCCAU CCGAU <u>GGACU</u> GCCCC GGAGG CCAUU GCCUU	N/A	N/A
14	1660	UUUGG GGAGA GGCG UACUG <u>GGACA</u> UGAGC AAUCA GGACG UGAUC	N/A	N/A
15	1699	GUGAU CAAUG CCAUU GAACA <u>GGACU</u> ACCGG CUGCC CCCGC CCCCA	N/A	N/A
16	1723	UACCG GCUGC CCCCG CCCCC <u>AGACU</u> GUCCC ACCUC CCUCC ACCAG	N/A	N/A
17	1756	UCCCU CCACC AGCUC AUGCU <u>GGACU</u> GUUGG CAGAA AGACC GGAAU	N/A	N/A
18	1771	AUGCU GGACU GUUGG CAGAA <u>AGACC</u> GGAAU GCCCG GCCCC GCUUC	N/A	N/A
19	1816	CCCCA GGUGG UCAGC GCCCU <u>GGACA</u> AGAUG AUCCG GAACC CCGCC	N/A	N/A
20	1891	GGGGC CUCAC ACCCU CUCCU <u>GGACC</u> AGCGG CAGCC UCACU ACUCA	N/A	N/A
21	2029	GUCAG CCAGA UCUCU GCUGA <u>GGACC</u> UGCUC CGAAU CGGAG UCACU	N/A	N/A
22	2060	GAAUC GGAGU CACUC UGGCG <u>GGACA</u> CCAGA AGAAA AUCUU GGCCA	N/A	N/A
23	2118	GAAGU CCCAG GCCAA GCCGG <u>GAACC</u> CCGGG UGGGA CAGGA GGACC	N/A	N/A
24	2130	CAAGC CGGGA ACCCC GGGUG <u>GGACA</u> GGAGG ACCGG CCCCG CAGUA	N/A	N/A
25	2138	GAACC CCGGG UGGGA CAGGA <u>GGACC</u> GGCCC CGCAG UACUG ACCUG	N/A	N/A
26	2156	GAGGA CCGGC CCCGC AGUAC <u>UGACC</u> UGCAG GAACU CCCCCA CCCCA	N/A	N/A

27	2166	CCCGC AGUAC UGACC UGCAG <u>GAACU</u> CCCCA CCCCA GGGAC ACCGC	N/A	N/A
28	2182	GCAGG AACUC CCCAC CCCAG <u>GGACA</u> CCGCC UCCCC AUUUU CCGGG	N/A	N/A
29	2216	CAUUU UCCGG GGCAG AGUGG <u>GGACU</u> CACAG AGGCC CCCAG CCCUG	N/A	N/A
30	2356	CACCC CCCAG CCACC UCGGG <u>GAACU</u> CCAGA CCAAG GGUGA GGGCG	N/A	N/A
31	2392	GUGAG GGCAG CUUUC CCUCA <u>GGACU</u> GGGUG UGACC AGAGG AAAAG	N/A	N/A

Score(binary)	Score(knn)	Score(spectrum)	Score(combined)	Decision
0.432	0.509	0.732	0.556	m^6A site (Low confidence)
0.542	0.752	0.56	0.56	m^6A site (Low confidence)
0.582	0.672	0.508	0.557	m^6A site (Low confidence)
0.449	0.593	0.698	0.555	m^6A site (Low confidence)
0.725	0.856	0.621	0.69	m^6A site (Very high confidence)
0.6	0.688	0.685	0.639	m^6A site (High confidence)
0.639	0.724	0.668	0.655	m^6A site (High confidence)
0.574	0.598	0.608	0.589	m^6A site (Moderate confidence)
0.709	0.819	0.672	0.7	m^6A site (Very high confidence)
0.651	0.556	0.693	0.663	m^6A site (High confidence)
0.552	0.579	0.63	0.584	m^6A site (Moderate confidence)
0.567	0.562	0.629	0.591	m^6A site (Moderate confidence)

0.738	0.775	0.614	0.69	m^6A site (Very high confidence)
0.609	0.743	0.638	0.627	m^6A site (High confidence)
0.7	0.684	0.623	0.668	m^6A site (High confidence)
0.649	0.848	0.681	0.671	m^6A site (High confidence)
0.778	0.802	0.763	0.773	m^6A site (Very high confidence)
0.484	0.645	0.739	0.594	m^6A site (Moderate confidence)
0.677	0.756	0.748	0.71	m^6A site (Very high confidence)
0.549	0.752	0.63	0.592	m^6A site (Moderate confidence)
0.561	0.618	0.67	0.607	m^6A site (Moderate confidence)
0.539	0.691	0.759	0.634	m^6A site (High confidence)
0.399	0.569	0.743	0.545	m^6A site (Low confidence)
0.636	0.83	0.726	0.681	m^6A site (Very high confidence)
0.572	0.794	0.744	0.652	m^6A site (High confidence)
0.462	0.694	0.729	0.58	m^6A site (Moderate confidence)

0.585	0.759	0.713	0.645	m^6A site (High confidence)
0.605	0.723	0.704	0.651	m^6A site (High confidence)
0.679	0.694	0.697	0.687	m^6A site (Very high confidence)
0.581	0.591	0.598	0.588	m^6A site (Moderate confidence)
0.749	0.889	0.503	0.658	m^6A site (High confidence)